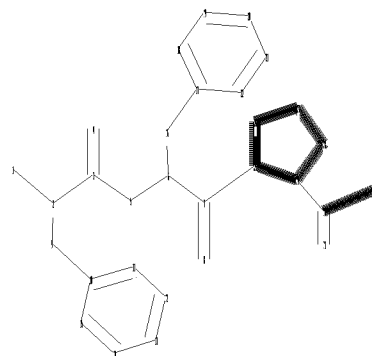
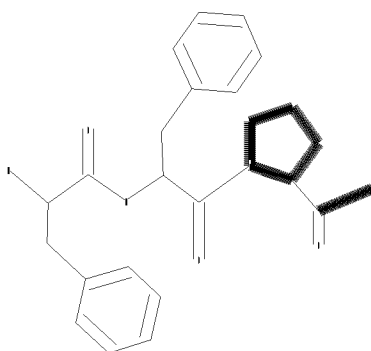


=>

Uploading C:\Program Files\Stnexp\Queries\86101.str



chain nodes :

1 2 3 4 5 6 7 8 9 16 30 31 32

ring nodes :

10 11 12 13 14 15 17 18 19 20 21 22 25 26 27 28 29

chain bonds :

1-2 2-3 2-9 3-4 3-5 5-6 6-7 6-16 7-8 7-25 9-10 16-17 29-30 30-31 30-32

ring bonds :

10-11 10-15 11-12 12-13 13-14 14-15 17-18 17-22 18-19 19-20 20-21 21-22
25-26 25-29 26-27 27-28 28-29

exact/norm bonds :

1-2 3-4 3-5 5-6 7-8 7-25 25-26 25-29 26-27 27-28 28-29 30-31 30-32

exact bonds :

2-3 2-9 6-7 6-16 9-10 16-17 29-30

normalized bonds :

10-11 10-15 11-12 12-13 13-14 14-15 17-18 17-22 18-19 19-20 20-21 21-22

G1:OH,MeO,EtO,n-PrO,i-PrO,n-BuO,i-BuO,s-BuO,t-BuO,PhO,NH2

Match level :

1:CLASS 2:CLASS 3:CLASS 4:CLASS 5:CLASS 6:CLASS 7:CLASS 8:CLASS 9:CLASS
10:Atom 11:Atom 12:Atom 13:Atom 14:Atom 15:Atom 16:CLASS 17:Atom 18:Atom
19:Atom 20:Atom
21:Atom 22:Atom 25:CLASS 26:Atom 27:Atom 28:Atom 29:Atom 30:CLASS 31:CLASS
32:CLASS

L7 STRUCTURE UPLOADED

=> s 17 sss full

FULL SEARCH INITIATED 13:13:12 FILE 'REGISTRY'

FULL SCREEN SEARCH COMPLETED - 316505 TO ITERATE

100.0% PROCESSED 316505 ITERATIONS

309 ANSWERS

SEARCH TIME: 00.00.05

L8 309 SEA SSS FUL L7

(FILE 'HOME' ENTERED AT 12:58:07 ON 13 APR 2009)

FILE 'REGISTRY' ENTERED AT 12:58:19 ON 13 APR 2009

L1 STRUCTURE UPLOADED

L2 25 S L1 SSS FUL

FILE 'HCAPLUS' ENTERED AT 12:59:20 ON 13 APR 2009

L3 1 S L2 AND ADRIAMYCIN

FILE 'STNGUIDE' ENTERED AT 12:59:35 ON 13 APR 2009

FILE 'REGISTRY' ENTERED AT 13:06:35 ON 13 APR 2009

L4 STRUCTURE UPLOADED

L5 6 S L4 SSS FULL

FILE 'HCAPLUS' ENTERED AT 13:08:00 ON 13 APR 2009

L6 2 S L5

FILE 'STNGUIDE' ENTERED AT 13:09:21 ON 13 APR 2009

FILE 'REGISTRY' ENTERED AT 13:11:33 ON 13 APR 2009

L7 STRUCTURE UPLOADED

L8 309 S L7 SSS FULL

FILE 'HCAPLUS' ENTERED AT 13:13:26 ON 13 APR 2009

=> s 18

L9 241 L8